

#21



SEQUENCE LISTING

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Singh, Lalji

<120> UNIVERSAL PRIMERS FOR WILDLIFE IDENTIFICATION

<130> U 013365-9

<140> 09/821,782
<141> 2001-03-29

<160> 255

<170> PatentIn version 3.1

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<223> Universal primer "mcb 398" for amplifying fragment of cytochrome b gene of animal species

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tactacacaa tcaaagatat cctgggcctt ctgtactaa tccttagcact catactactc 240
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tactacacaa tcaaagacat cctgggcctt ctagtactaa tcctaacact catactactc      240
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tactacacaa tcaaagacat cctgggcctt ctagtactaa tcctaacact catactactc      240
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tactacacaa tcaaagacat cctgggcctt ctgtactaa tcctaacact catactactc	240
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tactacacaa tcaaagacat cctgggcctt ctgtactaa tcctaacact catactactc	240
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<220>
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<223> DNA sequence generated from the known white tiger (*Panthera tigris tigris*) animal number 1 using primers mcb398 and mcb869

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<220>
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<223> DNA sequence generated from the known white tiger (*Panthera tigris tigris*) animal number 2 using primers mcb398 and mcb869

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<212> DNA
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<220>
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<223> DNA sequence generated from the known white tiger (Panthera tigris tigris) animal number 3 using primers mcb398 and mcb869

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<210> 15
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<212> DNA
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<220>
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tactacacaa tcaaagacat cctgggcctt ctgtactaa tcctaacact catactactc      240
gtcctattct caccagacct attagggac cccgataact acatccccgc caaccctcta      300
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<220>
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<223> DNA sequence generated from the known leopared (Panthera pardus) animal number 1 using primers mcb398 and mcb869

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<220>
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acaggatcta acaaccctc aggaatagta tccgactcag aaaaaattcc attccaccca      180
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gtcctattct caccagacct gtttaggagac cccgataact acatccctgc caaccctcta      300
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<223> DNA sequence generatd from the known leopared (Panthera pardus) a
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<220>
<221> misc_feature
<223> DNA sequence generated from the known leopared (Panthera pardus)
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acaggatcta acaaccctc aggaatagta tctgactcag aaaaaattcc attccaccca      180
tactacacaa tcaaagacat cctgggcctt ctgtactaa tccttagcact catactactc      240
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nimal number 3 using primers mcb398 and mcb869

<220>
<221> misc_feature
<223> DNA sequence generated from the known leopared (*Panthera pardus*) animal number 3 using primers mcb398 and mcb869

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acaggatcta acaaccctc aggaatagta tctgactcag aaaaaattcc attccaccca 180
tactacacaa tcaaagacat cctggccctt ctgtactaa tcttagcact catactactc 240
gtcctattct caccagacct gttgggagac cccgataact acatccccgc caaccctcta 300
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<210> 19
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<220>
<221> misc_feature
<223> DNA sequence generated from the known clouded leopard (*Neofelis n ebulosa*) animal number 1 using primers mcb398 and mcb869

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aaggatccaa taaccctca ggaatggtat ccgattcaga caaatcccc ttccacccgt 180
actatacat caaagatatac ctggcctcc tagttctaat tctagcgctc acactacttg 240
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<210> 20
<211> 327
<212> DNA
<213> gz22CL

<220>
<221> misc_feature
<223> DNA sequence generated from the known clouded leopard (*Neofelis n ebulosa*) animal number 2 using primers mcb398 and mcb869

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aaggatcaa taaccctca ggaatggat ccgattcaga caaaatccc ttccacccgt 180
actatacaat caaagatatac cttaggcctcc tagtctaat tctagcgctc acactacttg 240
ttctattctc cccagaccta cttaggagacc ctgacaatta cactccgccc aaccctctaa 300
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acaggatcta acaaccctc aggaatagta tctgacttag aaaaaatccc gttccaccca 180
tactacacaa tcaaagacat cctgggcctt ctagtactaa tcctaacact catactactc 240
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ttcatccttc catttatcat ctcagcccta gcagcagtcc acctcctatt cctccatgag 120
acaggatcta acaaccctc aggaatagta tctgacttag aaaaaatccc gttccaccca 180
tactacacaa tcaaagacat cctgggcctt ctagtactaa tcctaacact catactactc 240
gtcctattct caccagacct attagggac gccgataact acatccccgc caaccctctaa 300
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<223> DNA sequence generated from the known snow leopard (*Panthera uncia*) animal number 3 using primers mcb398 and mcb869

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tactacacaa tcaaagacat cctgggcctt ctgtactaa tcctaacact catactactc 240
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<222> (1)..(328)
<223> DNA sequence generated from the known asiatic lion (*Panthera leo persica*) animal number 1 using primers mcb398 and mcb869

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<220>
<221> misc_feature
<223> DNA sequence generated from the known asiatic lion (*Panthera leo persica*) animal number 1 using primers mcb398 and mcb869

persica) animal number 2 using primers mcb398 and mcb869

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acaggatcta ataacccttc aggaatggta tctgactcag ataaaattcc attccatcca 180
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agcacccctc cccatatcaa acctgaat 328

<210> 27
<211> 328
<212> DNA
<213> humsk

<220>
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<220>
<221> misc_feature
<223> DNA sequence generated from the known human (*Homo sapiens sapiens*) using primers mcb398 and mcb869

<220>
<221> misc_feature

<223> DNA sequence generated from the known human (*Homo sapiens sapiens*) using primers mcb398 and mcb869

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acgggatcaa acaacccctt aggaatcacc tcccattccg ataaaatcat cttccaccct 180
tactacacaa tcaaagacgc ctcggctta cttctttcc ttcttcctt aatgacatta 240
acactattct caccagacct cttaggcac ccagacaatt atacccttagc caacccctta 300
aacacccctc cccacatcaa gcccgaaat 328

<210> 28

<211> 328

<212> DNA

<213> chimss

<220>

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<223> DNA sequence gerated from the known chimpanzee (*pan troglodytes*) animal using primers mcb398 and mcb869

<220>

<221> misc_feature

<223> DNA sequence generated from the known chimpanzee (*pan troglodytes*) animal using primers mcb398 and mcb869

<220>

<221> misc_feature

<223> DNA sequence generated from the known chimpanzee (*pan troglodytes*) animal using primers mcb398 and mcb869

<400> 28

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acaggatcaa ataacccctt ggaaatcacc tcccactccg acaaaattac cttccacccc 180

tactacacaa tcaaagatat cttggctta ttcctttcc tccttatacct aatgacatta 240

acactattct caccagacct ctgggcgtat ccagacaact atacccttagc taacccctta 300

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328

<210> 29

<211> 472

<212> DNA

<213> Cervus nippon centralis

<400> 29

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caaccctaac ccgattttc gccttccact ttattcttcc atttatcatc gcagcacttg 180
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cgacgcaga caaaatcccc ttccatcctt actacaccat taaagatatac ttaggcattct 300
tacttctagt actcttccta atattactag tattattcgc accagacctg ctggagatc 360
cagacaacta tacccagca aatccactca acacacccccc tcacatcaaa cctgaatgtat 420
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<210> 30
<211> 472
<212> DNA
<213> Cervus nippon yesoensis

<400> 30
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caaccctaac ccgattttc gccttccact ttattcttcc atttatcatc gcagcacttg 180
ctatagtaca cttaactttc ctgcacgaga caggatccaa caacccaaca ggaatcccat 240
cgacgcaga caaaatcccc ttccatcctt actacaccat taaagatatac ttaggcattct 300
tacttctagt actcttccta atattactag tattattcgc accagacctg ctggagatc 360
cagacaacta tacccagca aatccactca acacacccccc tcacatcaaa cctgaatgtat 420
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<210> 31
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cagacgcaga caagatcccc ttccacccat actacaccat caaagatgtc ctaggggctc	300
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 tactactaat cctaacccta ctaatattaa ctctattgc acccgactta ctcggagacc 360
 cagacaacta caccccagca aacccactca gtaccccagc acacattaaa ccagagtgat 420
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 caatcgcca cctcattttc cttcacgaaa caggatccaa caacccaca ggcatccat 240
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 tattactaat cctaacccta ctaatgttaa ccctattcgc acctgacctg cttggagacc 360
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 <213> Balaenoptera physalus

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 caattgtcca ctttattttc cttcacgaaa caggatccaa caacccaca ggcatccat 240
 ccgacataga taaaatccca ttccacccctt accacacaat taaagacatt ctaggtgcc 300
 tattactaat cctaacccta ctaatactaa ccctattcgc acccgaccta cttggagacc 360
 cagacaacta taccccagca aacccactca gtaccccagc acacattaaa ccagaatgg 420

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ccaacataga	caaattcca	ttccacccct	actacacaat	taaagacatc	ctggcggtcc	300
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<212>	DNA					
<213>	<i>Cephalorhynchus commersonii</i>					
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caacactaac	acgcttttc	gccttccact	ttatcctccc	attcatcatc	acagcattag	180
cagccgtcca	cctactattc	ctacacgaaa	caggatccaa	caaccccaca	ggaatcccat	240
ccaacataga	cataatccca	ttccacccctt	attacacaat	taaagacatc	ctaggcgctt	300
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<213>	<i>Cephalorhynchus eutropia</i>					
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caacactaac	acgcttttc	gccttccact	ttatcctccc	attcatcatc	acagcattag	180

cagccgtcca cctactattc ctacacgaaa caggatccaa caaccccaca ggaatcccat	240
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tattccta at ctaacccta ct tagcactaa ccctattcgc ccctgaccta ctaggagacc	360
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tattccta at ctaacccta ct tagcactaa ccctattcgc ccctgaccta ctaggagacc	360
ctgataacta taccccgca aatccattaa gcaccccgac acacatcaa ccagaatggt	420
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<213> <i>Cephalorhynchus heavisidii</i>	
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<210> 71	
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<212> DNA	
<213> <i>cephalorhynchus hectori</i>	

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 ccaacataga cataatccca ttccaccctt attacacaat taaagacatc ttaggcgtt 300
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<210> 72
 <211> 472
 <212> DNA
 <213> *Lagenorhynchus australis*
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 tattccta at tcta at cc ta ct tagc actaa cc ctat tc gc cc t gac ct a c t a g g a c c 360
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<210> 73
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 <213> *Lagenorhynchus cruciger*
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<210> 74
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<212> DNA
<213> *Lagenorhynchus obscurus*

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<210> 75
<211> 472
<212> DNA
<213> *Lissodelphis borealis*

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<210> 76
<211> 472
<212> DNA
<213> *Lissodelphis peronii*

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<212>	DNA					
<213>	Globicephala macrorhynchus					
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<210>	78					
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<212>	DNA					
<213>	Globicephala melas					
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<210>	79					
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<213> *Feresa attenuata*

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<210> 80

<211> 472

<212> DNA

<213> *Peponocephala electra*

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ctaacaacta taccccagca aacccactaa gcacccctgc acacatcaaa ccagaatgtat 420
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<210> 81

<211> 472

<212> DNA

<213> *Grampus griseus*

<400> 81
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tactcctaat cctaacacta ctaacactaa ccctattcac ccctgaccta ctaggagacc 360

ctgataacta cactccagca aacccgctaa gcacccctgc acacatcaaa ccagaatgat	420
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<210> 82	
<211> 472	
<212> DNA	
<213> Pseudorca crassidens	
<400> 82	
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caacactaac acgtttttc actctccact ttatcctccc attcatcatt acagcactaa	180
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ccaacataga cataattcca ttccaccctt attacacaat taaagatatac cttaggcgccc	300
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<213> Lagenorhynchus acutus	
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caacactgac acgtttttc gccttccatt tcacccccc attcataatt acagcattag	180
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ctaacataga tataatcccg ttccaccctt attatacaat taaagatatac cttaggcgtt	300
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<210> 84	
<211> 472	
<212> DNA	
<213> Orcinus orca	
<400> 84	
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ccaacataga tataatccca ttccaccctt atcacacaaat taaagatacc cttaggcgcc 300
tactcttaat cctaaccctg ctagcactaa ccttattcgc ccctgaccta cttaggagacc 360
ctgacaacta taccccagca aatccactaa gcacccctgc acacatcaaa ccagaatgat 420
acttcctatt cgcatacgca atcctacgat cagttcccaa taaacttgga gg 472

<210> 85
<211> 472
<212> DNA
<213> *Orcaella brevirostris*

<400> 85
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caacactaac acgtttttc gccttccact ttatccttcc attcatcatc acagcactag 180
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ccaacataga cataatccca ttccaccctt atcatacatt taaagacatc cttaggcgcc 300
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ctgataacta tactccagca aatccactaa gcacccctgc acacatcaaa ccagaatgat 420
acttcctatt cgcatacgca atcctacgat caatcctaa taaactcggg gg 472

<210> 86
<211> 472
<212> DNA
<213> *Delphinus capensis*

<400> 86
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caacattaac acgcttttc gcttccact ttatccttcc attcatcatc acagcattag 180
cagccgttca cctgctattc ctacacgaaa caggatccaa taacccca ggaatcccat 240
ccaatataga cataatccca ttccaccctt attatacaat caaagatatc cttagtgcc 300
tactcctaattt ctaacccta ctagcactga ccctattcgc tccagaccta cttaggagacc 360
ctgataacta taccccagca aatccactaa gcacccctgc acatatcaaa ccagaatgat 420
actttctatt cgcatacgca atcttacgat caatcctaa taaacttgga gg 472

<210> 87
<211> 472

<212> DNA

<213> *Delphinus tropicalis*

<400> 87

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caacattaac acgcttttc gcttccact ttatcctccc attcatcatc acagcattag	180
cagccgttca cctgcttattc ctacacgaaa caggatccaa taacccaca ggaatcccat	240
ccaacataga cataatccca ttccaccctt attatacat caaagatatc ctaggtgcc	300
tactcctaattt cttaacctta ctagcactga ccctattcac tcccgaccta ctaggagacc	360
ctgataacta tacccagca aatccactaa gcacccctgc acatataaaa ccagaatgat	420
actttctattt cgcatatcgca atcttacgat caatccctaa taaacttggaa gg	472

<210> 88

<211> 472

<212> DNA

<213> *Delphinus delphis*

<400> 88

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caacattaac acgcttttc gcttccact ttatcctccc attcatcatc acagcactag	180
cagccgttca cctgcttattc ctacacgaaa caggatccaa taacccaca ggaatcccat	240
ccaatataca cataatccca ttccaccctt attatacat caaagatatc ctaggtgcct	300
tactcctaattt cttaacctta ctagcactaa ccctattcac tcccgaccta ctaggagacc	360
ctgataacta tacccagca aatccactaa gcacccctgc acacatataaaa ccagaatgat	420
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<210> 89

<211> 472

<212> DNA

<213> *Stenella clymene*

<400> 89

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caacattaac acgcttttc gcttccact ttatcctccc gttcatcatc acagcattag	180
cagccgttca cctgcttattc ctacacgaaa caggatccaa taacccaca ggaattccat	240
ccaatataca cataatccca ttccaccctt attatacat caaagatatc ctaggtgcct	300
tactcctaattt cttaacctta ctagcactaa ccctattcac ccccgaccta ctaggagacc	360

ctgacaacta taccccagca aatccactaa gcacccctgc acacatcaa ccagaatgat 420
actttctatt cgcatatgca atcttacgat caatccctaa taaacttgga gg 472

<210> 90
<211> 472
<212> DNA
<213> Stenella coeruleoalba

<400> 90
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caacattaac acgcttttc gcttccact ttatcctccc gttcattatc acagcattag 180
cagccgttca cctgctattc ctacacgaaa caggatccaa taacccaaca ggaattccat 240
ccaatataga cataattcca ttccaccctt attatacaat taaagatatc ctaggtgcct 300
tactccta at cttaacccta cttagcactaa ccctattcac ccccgaccta ctaggagacc 360
ctgataacta taccccagca aatccactaa gcacccctgc acacatcaa ccagaatgat 420
actttctatt cgcatatgca atcttacgat caatccctaa taaacttgga gg 472

<210> 91
<211> 472
<212> DNA
<213> Tursiops aduncus

<400> 91
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caacactaac acgcttttc gcttccact ttatcctccc gttcgtcatc acagcattag 180
cagccgttca cctgctattc ctacacgaaa caggatccaa taaccccaca ggaatccat 240
ccaatataga cataatccc ttccaccctt attatacaat caaagacatc ctaggtgcct 300
tactccta at cttaacccta cttagcactaa ccctattcac ccccgaccta ctaggaaacc 360
ctgataacta tatcccagca aatccactaa gtacccccgc acacatcaa ccagagtgat 420
actttctatt cgcatatgca atcttacgat caatccctaa taaacttgga gg 472

<210> 92
<211> 472
<212> DNA
<213> Stenella frontalis

<400> 92
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caacattaac acgcttttc gcttccact ttatcctccc gttcatcatc acagcattag	180
cagccgttca cctactattc ctacacgaaa caggatccaa taaccccaca ggaatcccat	240
ccaatataga cataatccca ttccaccctt attatacat caaagacatc cttaggcgcct	300
tactcctaatt ctaacccta cttagcactaa ccctattcac cccccgaccta cttaggagacc	360
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<210> 93	
<211> 472	
<212> DNA	
<213> Sousa chinensis	
<400> 93	
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caacattaac acgcttttc gcttccact ttatcttcc cttcatcatc acagcattag	180
tagccgttca cctgctattc ctacacgaaa caggatccaa taaccctaca ggaattccat	240
ccaacataga cataatccca ttccaccctt attatacat caaagacatc cttagtgccct	300
tactcctaatt ctaacccta cttagcactaa ccctattcac cccccgaccta cttaggagacc	360
ccgataacta tacccagca aatccactaa gcacccctgc acacatcaa ccagaatgat	420
atccctatt cgcatacgca atcttacgat caatccctaa taaaacttggaa gg	472

<210> 94	
<211> 472	
<212> DNA	
<213> Stenella longirostris	
<400> 94	
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caacattaac acgcttttc gcttccatt ttatcctccc attcatcatc acagcattag	180
cagccgtcca cctactattc ctacacgaaa caggatccaa taaccccaca ggaatcccat	240
ccaacataga cataatccca ttccaccctt attatacat caaagacatc cttagtggt	300
tactcttaat ctaacccta cttagcactaa ccctattcac ccctgactta cttaggagacc	360
ctgataacta tacccagca aatccactaa acacccctgc acacatcaa ccagaatgat	420
atccctatt cgcatacgca atcttacgat caatccctaa taaaacttggaa gg	472

<210> 95

<211> 472
 <212> DNA
 <213> *Tursiops truncatus*

<400> 95
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 cagccgttca cctactattc ctacacgaaa caggatccaa caaccccaca ggaatcccat 240
 ccaatataga cataatccca ttccaccctt attatacat caaagacatc ctaggcgcct 300
 tactcttaat cttAACCTTA cttagcattaa ccctattcgc ccccgaccta ctaggagacc 360
 ctgataacta cacccagca aacccactaa gcacccctgc acacatcaaa ccagaatgat 420
 actttctatt cgcatatcgca atcttacgat caatccctaa taagctcgga gg 472

<210> 96
 <211> 472
 <212> DNA
 <213> *Lagenorhynchus alborostris*

<400> 96
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 caacactaac acgcttcttc gctttccact ttatcctccc attcatcatc acagcactag 180
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 ccaacataga tataattcca ttccaccctt attacacaat caaagacatc ctaggcgcct 300
 tacttttaat cttAACCTTA cttagcactaa ccctatttac ccccgaccta ctaggagatc 360
 ccgataacta tacccagca aatccactaa gcactcctgc acacatcaaa ccagaatggt 420
 atttcctatt cgcatatcgca atcttacgat caatccctaa ccaaacttgga gg 472

<210> 97
 <211> 472
 <212> DNA
 <213> *Steno bredanensis*

<400> 97
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tacttttaat cctaacttta ctagcactaa ccctattcac ccccgaccta cttaggagacc	360
ccgacaacta taccccagca aatccactaa gcacccctgc acacatcaa ccagaatgg	420
attccttatt cgcatatcgca atcttacgt caatccccaa caaacttgga gg	472
<210> 98	
<211> 472	
<212> DNA	
<213> <i>Sotalia fluviatilis</i>	
<400> 98	
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caacactaac acgcttttc gccttccact ttatcctccc atttatcatc acagcattag	180
cagccgttca cctgctattc ctacacgaaa caggatccaa taatcccaca ggaatcccatt	240
ccaacataga tataattcca ttccacccctt attacacaaat caaagatatc ctaggcgcc	300
tactcctaatt cctgacccta cttagcactaa ccctattcac ccccgaccta cttaggagatc	360
ccgacaacta tactccagca aatccactta acacccctgc acacatcaa ccagaatggat	420
attccttatt cgcatatcgca atcttacgt caatccccaa taaacttgga gg	472
<210> 99	
<211> 472	
<212> DNA	
<213> <i>Delphinapterus leucas</i>	
<400> 99	
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caacactaac acgcttcttc accttccact ttatcctccc attcatcatt acagcgctag	180
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tactactaat cctaacccta ttaacagtaa ccctattcac acctgacccctc cttaggagacc	360
cagacaatta caccccttgc aacccactaa acaccccccgc acacatcaa ccagaatgg	420
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<210> 100	
<211> 472	
<212> DNA	
<213> <i>Monodon monoceros</i>	
<400> 100	
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tggccgtcca	cttattattc	ctacacgaaa	caggatccaa	caaccccaca	ggaatcccat	240
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ctgacaatta	tacccagca	aacccactaa	gcacccctgc	acacatcaaa	ccagaatgat	420
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<210> 101
<211> 472
<212> DNA
<213> Platanista gangetica

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ccgataacta	cacccagca	aacccgctta	atacccagc	acatatcaaa	ccagagtgtat	420	
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<210> 102
<211> 472
<212> DNA
<213> Platanista minor

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ccgataacta	cacccagca	aacccgctta	atacccagc	acatatcaaa	ccagagtgtat	420	
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<210> 103
 <211> 472
 <212> DNA
 <213> *Kogia breviceps*

<400> 103
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 ctgacaacta caccccagca aacccactaa gcaccccgac acacattaaa ccagaatgat 420
 atttcctatt tgcatatgcc atcctacgat ccattcctaa caaacttaggg gg 472

<210> 104
 <211> 472
 <212> DNA
 <213> *Kogia simus*

<400> 104
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 ctgatataga caaaatccca ttccacccct actacacaat caaagatatc ttaggcgcc 300
 tactactaat ctcagcacta ctcacactga ccctgttcgc acctgatcta ttaggagacc 360
 ccgacaacta taccccagca aacccactaa gcaccccgac acacattaaa ccagaatgat 420
 actttctatt cgcatatgcc attctacgat caattcctaa caaactggga gg 472

<210> 105
 <211> 472
 <212> DNA
 <213> *Physeter catodon*

<400> 105
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 caatagtaca tctccttattt ctccatgaaa caggatccaa caacccacca ggaattccct 240
 ccaacataga caaaatccca ttccacccct accacacaat caaagacacc ataggtgcc 300

tactactaat cctatcccta cttacactaa ccctgttcgc acccgacctg ctaggagatc 360
ctgacaacta caccccagca aatccactaa ataccccaac acacatcaaa ccagaatggt 420
attccttatt cgctacgat atcctacgat ctgtccccaa taaaacttagga gg 472

<210> 106
<211> 472
<212> DNA
<213> *Lipotes vexillifer*

<400> 106
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caacattaac ccgcttcttc gctctccatt tcattttcc atttattatt gtagcactaa 180
caaccgtcca cttactatcc ctccatgaaa caggatccaa caacccaata ggaattccat 240
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<210> 107
<211> 472
<212> DNA
<213> *Phocoena sinus*

<400> 107
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caacactaac acgcttcttc gccttccatt ttatccttcc atttattcatt acagcactaa 180
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<210> 108
<211> 472
<212> DNA
<213> *Berardius bairdii*

<400> 108
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caaccctaac acgattttc gccttccact tcatacttacc attcgttagta ttagcactag	180
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ccgactcaga caaaatcccg ctccacccat attatacat taaagatatc ctaggagccc	300
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<210> 141

<211> 472

<212> DNA

<213> Leptonychotes weddelli

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<210> 142

<211> 472

<212> DNA

<213> Mirounga leonina

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<210> 146
<211> 472
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<210> 147
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<213> *Ailurus fulgens*

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<210> 148
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<210> 155
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<212> DNA
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<210> 156

<211> 472
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<210> 157
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 <212> DNA
 <213> Galagooides demidoff

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<212> DNA
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ccacattaac	tcgattcttc	actttacact	tcctccttcc	attcataatc	ataggcctca	180
ccctaattcca	cctcaccttc	cttcacgaat	ccggctcaaa	caacccctta	ggcatcgtat	240
caaactgcga	taaaatccca	ttccacccct	acttttcctt	aaaagatatac	ctaggattca	300
cactcatact	acttccactc	ataaccctag	ccctattctc	accaaaccta	ctaggagacc	360
cagaaaaactt	caccccagca	aacccctag	tcacacctcc	tcatatcaag	ccagaatgat	420
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<210> 185						
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<212> DNA						
<213> <i>Grus antigone gillae</i>						

<400> 185
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 ccacattaac tcgatttttc actttacact tcctccttcc attcataatc ataggcctca 120
 ccctaattcca cctcaccttc cttcacgaat ccggctcaaa caacccctta ggcatcgtat
 caaactgcga taaaatccca ttccacccct actttcctt aaaagatatc ctaggattca 180
 cactcatact acttccactc ataaccctag ccctattctc accaaaccta ctaggagacc
 cagaaaaactt cacccagca aacccctag tcacacctcc tcataatcaag ccagaatgat 240
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<210> 186
 <211> 472
 <212> DNA
 <213> *Grus antigone sharpei*
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 caaactgcga taaaatccca ttccacccct actttcctt aaaagatatc ctaggattca 180
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 cagaaaaactt cacccagca aacccctag tcacacctcc ccataatcaag ccagaatgat 240
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 360
 420
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<210> 187
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 <212> DNA
 <213> *Grus leucogeranus*
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 cagaaaaactt cactccagca aacccctag taacaccccc acatattaaa ccagaatgat 240
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 360
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<213> Grus canadensis pratensis	
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cagaaaaactt cacccagca gacccctag tcacacccctcc ccatatcaaa ccagaatgat	420
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<212> DNA	
<213> Grus canadensis rowani	
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cagaaaaactt cacccagca aacccctag tcacacccctcc ccatatcaaa ccagaatgat	420
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ccacattaac ccgatttttc actttacact tcctcctccc attcataatt ataggcctca	180

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tactcatact	acttccactc	ataaccctag	ctctatttc	accaaactta	ctaggagacc	360
cagaaaactt	cacccagca	aacccctag	tcacacctcc	ccatatcaaa	ccagaatgat	420
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<210> 191
<211> 472
<212> DNA
<213> *Grus canadensis canadensis*

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caaactgcga	taaaatccca	ttccacccctt	attttcctt	aaaagatatac	ctagggttca	300
tactcatact	acttccactt	ataaccctag	ctctatttc	accaaactta	ctaggagacc	360
cagaaaactt	cacccagca	aacccctag	tcacacctcc	ccatatcaaa	ccagaatgat	420
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<210> 192
<211> 472
<212> DNA
<213> *Grus americana*

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cactcatatt	acttccactc	ataaccctag	ctctatttc	accaaactta	ctaggagacc	360
cagaaaactt	cacccagca	aacccctag	tgacacctcc	ccatattaag	ccggaatgat	420
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<210> 193
<211> 472
<212> DNA
<213> *Grus grus*

<400> 193
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 ccctaattcca cctcaccttc ctgcacgaat ccggctcaaa caacccctta ggcatcgtat 240
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<210> 194

<211> 472

<212> DNA

<213> Grus monacha

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 cagaaaactt cacccagca aaccctctag tcacacctcc tcataaaaa ccggaatgat 420
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<210> 195

<211> 472

<212> DNA

<213> Grus nigricollis

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 ccctaattcca cctcaccttc ctgcacgaat ccggctcaaa caacccctta ggcatcgtat 240
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 tattcatatt acttccactc ataaccctag ctctatttc accaaactta ctaggagacc 360
 cagaaaactt cacccagca aaccctctag tcacacctcc ccatattaag ccggaatgat 420

actttctatt tgcatacgct atcctacgtt caatccaaa caaactagga gg 472

<210> 196
<211> 472
<212> DNA
<213> Grus japonensis

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ccctaatacca tctcacttcc ctccacgaat ccggctcaaa caacccctta ggcatcgat 240
caaactgtga taaaatccca ttccacccct attttcctt aaaagatatc ttaggattta 300
cactcatatt acttccactc ataaccctag ccctattctc accaaactta ctaggagacc 360
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<210> 197
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<210> 198
<211> 472
<212> DNA
<213> Rhea americana

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cagaaaactt caccccagcc aacccctag ttacacccccc tcacatcaag ccagaatgat		420
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<210> 199
<211> 472
<212> DNA
<213> Anthracoceros albirostris

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caaccctgac acgattcttc gccctacact ttctcctccc gttcataatc gcaggcctag	180
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cagtaatact ctcctcccta acctccctag cccttcttc ccccaaccta cttaggagacc	360
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<210> 200
<211> 472
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<213> Falco femoralis

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caacactgac ccgattcttc gccctacact tcctcctacc attcctaatc gcaggcctca	180
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<210> 201
<211> 472
<212> DNA

<213> Falco verpertinus

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<210> 202

<211> 472

<212> DNA

<213> Falco peregrinus

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<210> 203

<211> 472

<212> DNA

<213> Falco sparverius

<400> 203
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<210> 204	
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<212> DNA	
<213> <i>Aythya americana</i>	
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<210> 205	
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<213> <i>Smithornis sharpei</i>	
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<213> <i>Vidua chalybeata</i>	
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cagaaaactt	cactccggcc	aaccccctaa	tcacaccacc	acatatcaa	cccgaaatgat	420
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<210> 207

<211> 472

<212> DNA

<213> Chrysemys picta

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<210> 208

<211> 472

<212> DNA

<213> Emys orbicularis

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caaccctaac	ccgatttttc	actttccatt	tcttactgcc	atttaccatt	ataggcctaa		180
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cagataactt	tacaccagct	aacccgctat	ccacccacc	acatattaag	ccagagtgat		420
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<210> 209

<211> 472

<212> DNA
<213> Chelonia mydas

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<213> Eumeces egregius

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cagattcaga taaaattcca ttccaccctt actatactat traagacatc ctaggaatcc      300
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<213> Addax nasomaculatus

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cagacaacta caccccgca aacccactta atacaccccc tcacatcaaa cccgaatgtat	420	
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<212> DNA
<213> Pseudois nayaur

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cagacaacta tacccagca aatccactca acacacccc tcacattaaa cctgaatgat 420
atttctatt tgcatacgca atcctacgat caattccaa caaactagga gg 472

<210> 239
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<213> *Rupicapra pyrenaica*

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ctaccctcac ccgattctt gccttcact tcattctcc attcatcatt gcagccttag 180
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cagataacta tacccagcg aacccactca acacacccc tcacatcaaa cccgaatgat 420
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<211> 472
<212> DNA
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cagatgcgga caaaatccca tttnacccct attataccat caaagacatt ctggcgcca      300
tactactaat cctcaccctc atactactag tactattnac acctgaccta ctggagacc      360
cagataatta cacccagcg aacccactca acacaccccc tcacattaaa cccgagtgtat      420
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<210> 241
<211> 472
<212> DNA
<213> Pantholops hodgsoni

<400> 241
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cagatgcaga caaaatccca tttcaccctt actataccat taaagacatc ctggcgcta      300
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<212> DNA
<213> Budorcas taxicolor taxicolor

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cagataatta	taccccgca	aatccactca	acacaccccc	tcacatcaa	cctgaatgat	420
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<211> 472
<212> DNA
<213> Ovis ammon

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cagacaacta	caccccgca	aacccactta	acactccccc	tcacatcaa	cctgaatgat	420
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cgAACACAGA caaaatcccc ttcnnnnnnn nnnnnnnnat taaagacatt ctgggtGCCA      300
tcctactaat cctcatcctc atgctgctag tactattcac gcctgactta ctTGGAGACC      360
cagacaacta cacCCAGCA aACCCACTTA acactcccc tcacatcaaa cctGAATGAT      420
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<212> DNA
<213> Capcornis crispus

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<212> DNA
<213> Ovibos moschatus

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tactactaat ccttaccctt atactactag tattattcac acccgaccta cttggagacc	360
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<212> DNA
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<210> 249
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<212> DNA

<213> Cephalophus maxwellii

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<210> 250

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<212> DNA

<213> Alces alces

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tccttctaat tctttcccta atgttattag tcctatttc acctgacctg ctggagacc	360	

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<211> 472	
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